SEQUENCE LISTING

æ	(1) GENERAL INFORMATION:
5	(i) APPLICANT:
	(A) NAME: University of Guelph (B) STREET: Gordon St.
10	(C) CITY: Guelph (D) STATE: Ontario
10	(E) COUNTRY: Canada
	(F) POSTAL CODE (ZIP): NIG 2W1
15	(ii) TITLE OF INVENTION: Monocot Transformation Using Agrobacterium
	(iii) NUMBER OF SEQUENCES: 4
	(iv) COMPUTER READABLE FORM:
20	(A) MEDIUM TYPE: Floppy disk (B) COMPUTER: IBM PC compatible
	(C) OPERATING SYSTEM: PC-DOS/MS-DOS(D) SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
	(v) CURRENT APPLICATION DATA:
25	APPLICATION NUMBER: FILING DATE:
	(2) INFORMATION FOR SEQ ID NO: 1:
30	
30	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs
	(B) TYPE: nucleic acid (C) STRANDEDNESS: single
35	(D) TOPOLOGY: linear
	(ii) MOLECULE TYPE: other nucleic acid
	(A) DESCRIPTION: /desc = "primer 1"
40	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
	CCGTCTGCGG GAGCGCTATC C 21
45	(2) INFORMATION FOR SEQ ID NO: 2:
	(i) SEQUENCE CHARACTERISTICS:
* 0	(A) LENGTH: 21 base pairs
50	(B) TYPE: nucleic acid (C) STRANDEDNESS: single
	(D) TOPOLOGY: linear
55	<pre>(ii) MOLECULE TYPE: other nucleic acid</pre>
JJ	(A) DESCRIPTION: / desc = "primer 2"

5	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:	
J	CATCGCAAGA CCGGCAACAG G	21
	(2) INFORMATION FOR SEQ ID NO: 3:	
10	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 18 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
13	<pre>(ii) MOLECULE TYPE: other nucleic acid</pre>	
20		
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:	
25	AGAAACCAAA GGGTCCTG	18
20	(2) INFORMATION FOR SEQ ID NO: 4:	
30	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
35	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "primer 4"</pre>	
40	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:	
	02.002.02.000 2.00mm2.00	10